SEQUENCE LISTING

<110>   Chano, Tokuhiro
         Okabe, Hidetoshi
         Ikegawa, Shiro

<120>   RB1 gene induced protein (RB1CC1) and gene

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<140>   US 10/516,558
<141>   2004-11-30

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<151>   2003-01-30

<150>   JP P2002-161400
<151>   2002-06-03

<150>   JP P2002-214978
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Gln Ser Lys Tyr Lys Ile Ala Ile Gln His Gln Val Leu Val Val Asn
        35   40   45

Gly Gly Glu Cys Met Ala Ala Asp Arg Arg Val Cys Thr Tyr Ser Ala
    50   55   60

Gly Thr Asp Thr Asn Pro Ile Phe Leu Phe Asn Lys Glu Met Ile Leu
65   70   75   80

Cys Asp Arg Pro Pro Ala Ile Pro Lys Thr Thr Phe Ser Thr Glu Asn
                85   90   95

Asp Met Glu Ile Lys Val Glu Glu Ser Leu Met Met Pro Ala Val Phe
              100   105   110

His Thr Val Ala Ser Arg Thr Gln Leu Ala Leu Glu Met Tyr Glu Val
              115   120   125

Ala Lys Lys Leu Cys Ser Phe Cys Glu Gly Leu Val His Asp Glu His
    130   135   140

Leu Gln His Gln Gly Trp Ala Ala Ile Met Ala Asn Leu Glu Asp Cys
145   150   155   160

Ser Asn Ser Tyr Gln Lys Leu Leu Phe Lys Phe Glu Ser Ile Tyr Ser
              165   170   175

Asn Tyr Leu Gln Ser Ile Glu Asp Ile Lys Leu Lys Leu Thr His Leu
              180   185   190

Gly Thr Ala Val Ser Val Met Ala Lys Ile Pro Leu Leu Glu Cys Leu
              195   200   205

Thr Arg His Ser Tyr Arg Glu Cys Leu Gly Arg Leu Asp Ser Leu Pro
    210   215   220

Glu His Glu Asp Ser Glu Lys Ala Glu Thr Lys Arg Ser Thr Glu Leu
225   230   235   240

Val Leu Ser Pro Asp Met Pro Arg Thr Thr Asn Glu Ser Leu Leu Thr
              245   250   255

Ser Phe Pro Lys Ser Val Glu His Val Ser Pro Asp Thr Ala Asp Ala
              260   265   270

Glu Ser Gly Lys Glu Ile Arg Glu Ser Cys Gln Ser Thr Val His Gln
    275   280   285

Gln Asp Glu Thr Thr Ile Asp Thr Lys Asp Gly Asp Leu Pro Phe Phe
    290   295   300

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Asn Val Ser Leu Leu Asp Trp Ile Asn Val Gln Asp Arg Pro Asn Asp
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Val Glu Ser Leu Val Arg Lys Cys Phe Asp Ser Met Ser Arg Leu Asp
                325             330             335


Pro Arg Ile Ile Arg Pro Phe Ile Ala Glu Cys Arg Gln Thr Ile Ala
            340             345             350


Lys Leu Asp Asn Gln Asn Met Lys Ala Ile Lys Gly Leu Glu Asp Arg
        355             360             365


Leu Tyr Ala Leu Asp Gln Met Ile Ala Ser Cys Gly Arg Leu Val Asn
    370             375             380


Glu Gln Lys Glu Leu Ala Gln Gly Phe Leu Ala Asn Gln Lys Arg Ala
385             390             395             400


Glu Asn Leu Lys Asp Ala Ser Val Leu Pro Asp Leu Cys Leu Ser His
            405             410             415


Ala Asn Gln Leu Met Ile Met Leu Gln Asn His Arg Lys Leu Leu Asp
            420             425             430


Ile Lys Gln Lys Cys Thr Thr Ala Lys Gln Glu Leu Ala Asn Asn Leu
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His Val Arg Leu Lys Trp Cys Cys Phe Val Met Leu His Ala Asp Gln
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Asp Gly Glu Lys Leu Gln Ala Leu Leu Arg Leu Val Ile Glu Leu Leu
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Glu Arg Val Lys Ile Val Glu Ala Leu Ser Thr Val Pro Gln Met Tyr
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Cys Leu Ala Val Val Glu Val Val Arg Arg Lys Met Phe Ile Lys His
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Tyr Arg Glu Trp Ala Gly Ala Leu Val Lys Asp Gly Lys Arg Leu Tyr
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Ser Phe Cys Thr Gln Lys Pro Arg Lys Phe Asp Cys Glu Leu Pro Asp
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Ile Ser Leu Lys Asp Leu Gln Phe Leu Gln Ser Phe Cys Pro Ser Glu
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Val Gln Pro Phe Leu Arg Val Pro Leu Leu Cys Asp Phe Glu Pro Leu
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His Gln His Val Leu Ala Leu His Asn Leu Val Lys Ala Ala Gln Ser
   610   615   620

Leu Asp Glu Met Ser Gln Thr Ile Thr Asp Leu Leu Ser Glu Gln Lys
625   630   635   640

Ala Ser Val Ser Gln Thr Ser Pro Gln Ser Ala Ser Ser Pro Arg Met
                645   650   655

Glu Ser Thr Ala Gly Ile Thr Thr Thr Ser Pro Arg Thr Pro Pro
                660   665   670

Pro Leu Thr Val Gln Asp Pro Leu Cys Pro Ala Val Cys Pro Leu Glu
                675   680   685

Glu Leu Ser Pro Asp Ser Ile Asp Ala His Thr Phe Asp Phe Glu Thr
   690   695   700

Ile Pro His Pro Asn Ile Glu Gln Thr Ile His Gln Val Ser Leu Asp
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Leu Asp Ser Leu Ala Glu Ser Pro Glu Ser Asp Phe Met Ser Ala Val
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Asn Glu Phe Val Ile Glu Glu Asn Leu Ser Ser Pro Asn Pro Ile Ser
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Asp Pro Gln Ser Pro Glu Met Met Val Glu Ser Leu Tyr Ser Ser Val
     755   760   765

Ile Asn Ala Ile Asp Ser Arg Arg Met Gln Asp Thr Asn Val Cys Gly
     770   775   780

Lys Glu Asp Phe Gly Asp His Thr Ser Leu Asn Val Gln Leu Glu Arg
785   790   795   800

Cys Arg Val Val Ala Gln Asp Ser His Phe Ser Ile Gln Thr Ile Lys
            805   810   815

Glu Asp Leu Cys His Phe Arg Thr Phe Val Gln Lys Glu Gln Cys Asp
            820   825   830

Phe Ser Asn Ser Leu Lys Cys Thr Ala Val Glu Ile Arg Asn Ile Ile
        835   840   845

Glu Lys Val Lys Cys Ser Leu Glu Ile Thr Leu Lys Glu Lys His Gln
     850   855   860

Lys Glu Leu Leu Ser Leu Lys Asn Glu Tyr Glu Gly Lys Leu Asp Gly
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Leu Ile Lys Glu Thr Glu Glu Asn Glu Asn Lys Ile Lys Lys Leu Lys
            885   890   895

Gly Glu Leu Val Cys Leu Glu Glu Val Leu Gln Asn Lys Asp Asn Glu
            900   905   910

Phe Ala Leu Val Lys His Glu Lys Glu Ala Val Ile Cys Leu Gln Asn
        915   920   925

Glu Lys Asp Gln Lys Leu Leu Glu Met Glu Asn Ile Met His Ser Gln
     930   935   940

Asn Cys Glu Ile Lys Glu Leu Lys Gln Ser Arg Glu Ile Val Leu Glu
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Asp Leu Lys Lys Leu His Val Glu Asn Asp Glu Lys Leu Gln Leu Leu
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Arg Ala Glu Leu Gln Ser Leu Glu Gln Ser His Leu Lys Glu Leu Glu
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Asp Thr Leu Gln Val Arg His Ile Gln Glu Phe Glu Lys Val Met Thr
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Asp His Arg Val Ser Leu Glu Glu Leu Lys Lys Glu Asn Gln Gln
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Ile Ile Asn Gln Ile Gln Glu Ser His Ala Glu Ile Ile Gln Glu
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Lys Glu Lys Gln Leu Gln Glu Leu Lys Leu Lys Val Ser Asp Leu
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Ser Asp Thr Arg Cys Lys Leu Glu Val Glu Leu Ala Leu Lys Glu
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Ala Glu Thr Asp Glu Ile Lys Ile Leu Leu Glu Glu Ser Arg Ala
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Gln Gln Lys Glu Thr Leu Lys Ser Leu Leu Glu Gln Glu Thr Glu
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Asn Asn Glu Asn Tyr Gln Val Gly Leu Ala Glu Leu Arg Thr Leu
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Met Thr Ile Glu Lys Asp Gln Arg Ile Ser Glu Leu Ile Ser Arg
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His Glu Glu Glu Ser Asn Ile Leu Lys Ala Glu Leu Asn Lys Val
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Thr Ser Leu His Asn Gln Ala Phe Glu Ile Glu Lys Asn Leu Lys
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Glu Gln Ile Ile Glu Leu Gln Ser Lys Leu Asp Ser Glu Leu Ser
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Leu Val  Ser Ser Gln Glu Gln  Asp Arg Glu Gln Leu  Ile Gln Lys
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Leu Asn  Cys Glu Lys Asp Glu  Ala Ile Gln Thr Ala  Leu Lys Glu
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Phe Lys  Leu Glu Arg Glu Val  Val Glu Lys Glu Leu  Leu Glu Lys
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Val Lys  His Leu Glu Asn Gln  Ile Ala Lys Ser Pro  Ala Ile Asp
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Ser Thr  Arg Gly Asp Ser Ser  Ser Leu Val Ala Glu  Leu Gln Glu
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Leu Ile  Ala Glu Gln Gln Thr  Asn Phe Asn Thr Val  Leu Thr Arg
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Glu Lys  Met Arg Lys Glu Asn  Ile Ile Asn Asp Leu  Ser Asp Lys
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Leu Lys  Ser Thr Met Gln Gln  Gln Glu Arg Asp Lys  Asp Leu Ile
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Glu Ser  Leu Ser Glu Asp Arg  Ala Arg Leu Leu Glu  Glu Lys Lys
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Ser Val  Gln Glu Asn Ile His  Met Leu Ser Glu Glu  Lys Gln Arg
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Ile Met  Leu Leu Glu Arg Thr  Leu Gln Leu Lys Glu  Glu Glu Asn
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Lys Arg  Leu Asn Gln Arg Leu  Met Ser Gln Ser Met  Ser Ser Val
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Ser Ser  Arg His Ser Glu Lys  Ile Ala Ile Arg Asp  Phe Gln Val
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Gly Asp  Leu Val Leu Ile Ile  Leu Asp Glu Arg His  Asp Asn Tyr
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Val Leu  Phe Thr Val Ser Pro  Thr Leu Tyr Phe Leu  His Ser Glu
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Tyr Cys  Gln Ala Lys Lys Ala  Gln Asn Arg Phe Lys  Val Pro Leu
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            35   40   45

Gly Gly Glu Cys Met Ala Ala Asp Arg Arg Val Cys Thr Tyr Ser Ala
        50   55   60

Gly Thr Asp Thr Asn Pro Ile Phe Leu Phe Asn Lys Glu Met Ile Leu
65   70   75   80

Cys Asp Arg Ala Pro Ala Ile Pro Lys Ala Thr Phe Ser Thr Glu Asn
                85   90   95

Asp Met Glu Ile Lys Val Glu Glu Ser Leu Met Met Pro Ala Val Phe
            100   105   110

His Thr Val Ala Ser Arg Thr Gln Leu Ala Val Glu Met Tyr Asp Val
            115   120   125

Ala Lys Lys Leu Cys Ser Phe Cys Glu Gly Leu Val His Asp Glu His
        130   135   140

Leu Gln His Gln Gly Trp Ala Ala Ile Met Ala Asn Leu Glu Asp Cys
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Ser Asn Ser Tyr Gln Lys Leu Leu Phe Lys Phe Glu Ser Ile Tyr Ser
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Asp Tyr Leu Gln Ser Ile Glu Asp Ile Lys Leu Lys Leu Thr His Leu
            180   185   190

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Glu His Glu Gly Ser Glu Lys Ala Glu Met Lys Arg Ser Thr Glu Leu
225                 230                 235                 240


Val Leu Ser Pro Asp Met Pro Arg Thr Thr Asn Thr Ser Leu Val Thr
                245                 250                 255


Ser Phe His Lys Ser Met Glu His Val Ala Pro Asp Pro Thr Gly Thr
            260                 265                 270


Glu Arg Gly Lys Glu Leu Arg Glu Ser Cys Gln Ser Thr Val Gln Gln
        275                 280                 285


Glu Glu Ala Ser Val Asp Ala Lys Asp Ser Asp Leu Pro Phe Phe Asn
    290                 295                 300


Val Ser Leu Leu Asp Trp Ile Asn Val Gln Asp Arg Pro Asn Asp Val
305                 310                 315                 320


Glu Ser Leu Val Arg Lys Cys Phe Asp Ser Met Ser Arg Leu Asp Pro
            325                 330                 335


Lys Ile Ile Gln Pro Phe Met Leu Glu Cys His Gln Thr Ile Ala Lys
            340                 345                 350


Leu Asp Asn Gln Asn Met Lys Ala Ile Lys Gly Leu Glu Asp Arg Leu
        355                 360                 365


Tyr Ala Leu Asp Gln Met Ile Ala Ser Cys Ser Arg Leu Val Asn Glu
    370                 375                 380


Gln Lys Glu Leu Ala Gln Gly Phe Leu Ala Asn Gln Met Arg Ala Glu
385                 390                 395                 400


Asn Leu Lys Asp Ala Ser Val Leu Pro Asp Leu Cys Leu Ser His Ala
                405                 410                 415
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Asn Gln Leu Met Ile Met Leu Gln Asn His Arg Lys Leu Leu Asp Ile
        420   425   430

Lys Gln Lys Cys Thr Thr Ala Lys Gln Glu Leu Ala Asn Asn Leu His
        435   440   445

Val Arg Leu Lys Trp Cys Cys Phe Val Met Leu His Ala Asp Gln Asp
    450   455   460

Gly Glu Lys Leu Gln Ala Leu Leu Arg Leu Val Ile Glu Leu Leu Glu
465   470   475   480

Arg Val Arg Ile Val Glu Ala Leu Ser Thr Val Pro Gln Met Tyr Cys
            485   490   495

Leu Ala Val Val Glu Val Val Arg Arg Lys Met Phe Ile Lys His Tyr
        500   505   510

Arg Glu Trp Ala Gly Ala Leu Val Lys Asp Gly Lys Gln Leu Tyr Glu
        515   520   525

Ala Glu Lys Ser Lys Arg Glu Ser Phe Gly Lys Leu Phe Arg Lys Ser
    530   535   540

Phe Leu Arg Asn Arg Leu Phe Lys Gly Leu Asp Ser Trp Pro Ser Ser
545   550   555   560

Phe Cys Thr Gln Lys Pro Arg Lys Phe Asp Cys Glu Leu Pro Asp Ile
            565   570   575

Ser Leu Lys Asp Leu Gln Phe Leu Gln Ser Phe Cys Pro Ser Glu Val
        580   585   590

Gln Pro Phe Leu Arg Val Pro Leu Leu Cys Asp Phe Glu Pro Leu His
        595   600   605

Gln His Val Leu Ala Leu His Asn Leu Val Lys Ala Ala Gln Ser Leu
    610   615   620

Asp Glu Met Ser Gln Thr Ile Thr Asp Leu Leu Asn Glu Gln Lys Val
625   630   635   640

Ser Thr Ser Gln Ala Ser Pro Gln Ser Ala Ala Ser Pro Arg Ile Glu
           645   650   655

Ser Thr Thr Gly Ile Thr Thr Thr Thr Ser Pro Lys Thr Pro Pro Pro
           660   665   670

Leu Thr Val Gln Asp Thr Leu Cys Pro Ala Val Cys Pro Leu Glu Glu
           675   680   685

Leu Ser Pro Asp Ser Ile Asp Ala His Thr Phe Asp Phe Glu Thr Ile
    690   695   700

Ser His Pro Asn Thr Glu Gln Pro Val His Gln Ala Ser Ile Asp Leu
705   710   715   720

Asp Ser Leu Ala Glu Ser Pro Glu Ser Asp Phe Met Ser Ala Val Asn
               725   730   735

Glu Phe Val Ile Glu Glu Asn Leu Ser Ser Pro Asn Pro Ile Ser Asp
               740   745   750

Pro Gln Ser Pro Glu Met Met Val Glu Ser Leu Tyr Ser Ser Val Ile
           755   760   765

Asn Ala Ile Asp Ser Arg Arg Met Gln Asp Thr Ser Thr Arg Gly Asn
    770   775   780

Glu Gly Phe Gly Asp Arg Ala Ala Leu His Val Gln Leu Glu Lys Cys
785   790   795   800

Arg Ala Ala Ala Gln Asp Ser His Thr Ser Ile Gln Thr Ile Lys Asp
               805   810   815

Asp Leu Cys His Phe Arg Thr Phe Val Gln Lys Glu Gln Cys Asp Leu
           820   825   830

Ala Asn Tyr Leu Lys Cys Thr Ala Val Glu Ile Arg Asn Ile Ile Glu
           835   840   845

Lys Val Lys Cys Ser Leu Glu Ile Thr Leu Lys Glu Lys His Gln Gln
    850   855   860

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Glu Leu Gln Ser Leu Lys Ile Glu Tyr Glu Cys Lys Leu Asp Ala Leu
865   870   875   880

Val Lys Asp Ser Glu Glu Asn Val Asn Lys Ile Leu Lys Leu Lys Glu
              885   890   895

Asn Leu Val Ser Leu Glu Glu Ala Leu Gln Asn Lys Asp Asn Glu Phe
          900   905   910

Thr Ser Ile Lys His Glu Lys Asp Ala Ile Val Cys Val Gln Gln Glu
          915   920   925

Lys Asp Gln Lys Leu Leu Glu Met Glu Lys Ile Met His Thr Gln His
    930   935   940

Cys Glu Ile Lys Glu Leu Lys Gln Ser Arg Glu Met Ala Leu Glu Asp
945   950   955   960

Leu Lys Lys Leu His Asp Glu Lys Ile Glu Ser Leu Arg Ala Glu Phe
              965   970   975

Gln Cys Leu Glu Glu Asn His Leu Lys Glu Leu Glu Asp Thr Leu His
          980   985   990

Ile Arg His Thr Gln Glu Phe Glu Lys Val Met Thr Asp His Asn Met
          995   1000   1005

Ser Leu Glu Lys Leu Lys Lys Glu Asn Gln Gln Arg Ile Asp Gln
    1010   1015   1020

Met Leu Glu Ser His Ala Ser Thr Ile Gln Glu Lys Glu Gln Gln
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Leu Gln Glu Leu Lys Leu Lys Val Ser Asp Leu Ser Asp Met Arg
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Cys Lys Leu Glu Val Glu Leu Ala Leu Lys Glu Ala Glu Thr Asp
    1055   1060   1065

Glu Ile Lys Ile Leu Leu Glu Glu Ser Arg Thr Gln Gln Lys Glu
    1070   1075   1080

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Tyr Gln  Val Gly Leu Ser Glu  Leu Arg Ala Leu Met  Thr Ile Glu
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Leu Ile  Gln Asn Leu Glu Lys  Asp Lys Glu Arg Leu  Val Lys Asn
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His Glu  Gln Asp Lys Glu His  Leu Ile Gln Glu Leu  Asn Phe Glu
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Lys Asn  Lys Ala Val Gln Thr  Ala Leu Asp Glu Phe  Lys Val Glu
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Arg Glu  Leu Val Glu Lys Glu  Leu Leu Glu Lys Val  Lys His Leu
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Glu Asn  Gln Ile Ala Lys Thr  Pro Ala Phe Glu Ser  Ala Arg Glu
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Asp Ser  Ser Ser Leu Val Ala  Glu Leu Gln Glu Lys  Leu Gln Glu
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Gln Gln  Thr Asn Phe Asn Thr  Val Leu Thr Arg Glu  Lys Met Arg
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Lys Glu  Asn Ile Ile Asn Asp  Leu Ser Asp Lys Leu  Lys Ser Thr
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Met Gln  Gln Gln Glu Arg Asp  Lys Asp Leu Ile Glu  Ser Leu Ser
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Glu Asp  Arg Ala Arg Leu Leu  Glu Glu Lys Lys Gln  Leu Glu Glu
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Glu Val  Ser Lys Leu Arg Thr  Ser Ser Phe Leu Ser  Ser Ala Pro
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Val Ala  Ala Ala Pro Glu Leu  Tyr Gly Ala Cys Ala  Pro Glu Leu
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Pro Gly  Glu Pro Glu Arg Ser  Val Met Glu Thr Ala  Asp Glu Gly
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Arg Leu  Asp Ser Ala Met Glu  Thr Ser Met Met Ser  Val Gln Glu
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Asn Met  Leu Ser Glu Glu Lys  Gln Arg Ile Met Leu  Leu Glu Arg
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Thr Leu  Gln Leu Lys Glu Glu  Glu Asn Lys Arg Leu  Asn Gln Arg
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Leu Met  Ser Gln Ser Leu Ser  Ser Val Ser Ser Arg  His Ser Glu
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Lys Ile  Ala Ile Arg Asp Phe  Gln Val Gly Asp Leu  Val Leu Ile
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Ile Leu  Asp Glu Arg His Asp  Asn Tyr Val Leu Phe  Thr Val Ser
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Pro Thr  Leu Tyr Phe Leu His  Ser Glu Ser Leu Pro  Ala Leu Asp
    1520              1525              1530

Leu Lys  Pro Gly Glu Gly Ala  Ser Gly Ala Ser Arg  Arg Pro Trp
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Val Leu  Gly Lys Val Met Glu  Lys Glu Tyr Cys Gln  Ala Lys Lys
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cccatccaaa catagaacag actattcacc aagtttcttt agacttggat tcattagcag    2700

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-19-

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-21-

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<210>  25
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<400> 26
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<400> 27
cctccctgcc tcctagagtt   20

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gagtgaaagc cgtatcatgg a                                              21


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aatgcggacc aaactacttc a                                              21


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tgattgctgg gaagtgtgaa                                              20


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<400>  36
gagaattgtt tgaggtttgt tgc                                          23


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<400>  37
ttgctcaatg gcaacttctc                                              20


<210>  38
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<400>  38
gggtgaggta taagctcaca gaa                                          23


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<400> 39
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caaaggattc cctttttgat ttt   23

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gctaatcaga agagagctga aaactt   26

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taagcatgcc attcaaagca   20

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<400>  44
tttctttata cagggaagtc ttt                                          23


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<400>  45
ttttcagaat gccttgaaga t                                            21


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aggataccat catcctagac gaa                                          23


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<400>  47
cttaccaccc tcacctggtt                                              20


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<210> 50
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<400> 50
aaaatatagg atacaaatcc aatgaca   27

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<400>  53
ggttctctga gtttcaccag taa                                        23


<210>  54
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<400>  54
atcaacactg gagggtgagg                                            20


<210>  55
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<400>  55
atcatcatga agttatatgt gtttctgg                                   28


<210>  56
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gatgcctgca gttttccaca                                            20


<210>  57
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<220>
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<400>  57
acgtggcaaa gaacttaggg                                            20
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<400>  58
gcttcttctt gctggacagt                                               20


<210>  59
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<400>  59
ccttggacca gatgattgct                                               20



<210>  60
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gggtcccctt actttgtgac t                                             21



<210>  61
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catccaaact ttgtgctgct                                               20



<210>  62
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tgctgcacaa gactctcaca                                          20


<210>  63
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<400>  63
ggcacagatc gtccttgatg g                                        21


<210>  64
<212>  DNA
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<220>
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<400>  64
tgaacttgca ctaaaggaag ca                                       22


<210>  65
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<400>  65
cagcatttcc ttctgctgtg                                          20


<210>  66
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<400>  66
tgagtgctct tgaaaaacag aaag                                     24


<210>  67
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<220>
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<400>   67
ttgcggaact tcaagagaaa c   21

<210>   68
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<400>   68
ctggaacaac ttgaagaaca agag   24

<210>   69
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acgagctcgg tcctcagaga   20

<210>   70
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tcaggtggga gatttggttc   20

<210>   71
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<400>   71
tgccgctcat ctaggatgat   20

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<210>  72
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<400>  72
cagcactgga ggacaaatca                                           20


<210>  73
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<400>  73
agtcacaagc cacagtgcag                                           20


<210>  74
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<400>  74
tgctttgaat ggcatgctta                                           20


<210>  75
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<400>  75
cctcaccctc cagtgttgat                                           20


<210>  76
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cctccgcacc atcttctg                                                      18


<210>  77
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<400>  77
cagggtcccc gtaggact                                                      18


<210>  78
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<400>  78
tcaggtggga gatttggttc                                                    20


<210>  79
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<220>
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<400>  79
ccagcatttg tcctcttgtg                                                    20


<210>  80
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<212>  DNA
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<400>  80
ttttgagttt gcctcagaag a                                                  21


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tcggaattca tggttgacct   20

<210>   82
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<220>
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<400>   82
tttcccagaa atcacgcaat   20

<210>   83
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<220>
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<400>   83
tctttgttca gtgccagctt t   21

<210>   84
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<220>
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<400>   84
accaggtggg tttgtcagag   20

<210>   85
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<400>   85
cttggcgatg caaagatgta   20

<210>   86
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acactggagg gtgaggtgtc                                              20


<210>   87
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<400>   87
gtgtcaaatg tcagcgtggt                                              20


<210>   88
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<400>   88
gacggttgtg tcggttgg                                                18


<210>   89
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<400>   89
ttggcaactg agtggcatta                                              20


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tgccactcag ttgccaagta                                             20
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<400> 91
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<210> 92
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<400> 92
cccagcgctg taagtacaca   20

<210> 93
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<400> 93
taagcatgcc attcaaagca   20

<210> 94
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<400> 94
caggtgcacg gtcacataag   20

<210> 95
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gcagttttcc acactgttgc                                                    20


<210>  96
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<400>  96
ctccagattg gccatgattg                                                    20


<210>  97
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<220>
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<400>  97
gccaatctgg aggactgttc                                                    20


<210>  98
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<212>  DNA
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<220>
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<400>  98
agaatccggt cttcccaaac                                                    20


<210>  99
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cccaatgatg tggaatctct g                                                  21

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<210> 101
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<210> 102
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<220>
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ggatgcatct gtgttacctg a   21

<210> 104
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<220>
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<400> 104
gcctgcagtt tttctccatc   20

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<400>  105
gctccgcctt gtaatagagc                                           20


<210>  106
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<400>  106
ccaaaggatt ccctttttga                                           20


<210>  107
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gggctggtgc tttagtcaaa                                           20


<210>  108
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<400>  108
aaaatgagga aggccaggag                                           20


<210>  109
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<220>
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actcctggcc ttcctcattt                                              20


<210>  110
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<400>  110
tgaggaatgg ctgcacttct                                              20


<210>  111
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<400>  111
gcctcgaaaa tttgactgtg a                                            21


<210>  112
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<400>  112
tccaaacttt gtgctgcttt t                                            21


<210>  113
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<400>  113
aaaagcagca caaagtttgg a                                            21


<210>  114
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<400>  114
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<210>  115
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<400>  115
ccacgagcaa gacaaagaac                                       20


<210>  116
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<220>
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<400>  116
tccgcaacta agcttgaaga a                                     21


<210>  117
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<400>  117
tgaggaaatg caaaatgtca                                       20


<210>  118
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<400>  118
gctcttgctg ctgcattgta                                       20
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<210> 119
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<400> 119
aaagtacaat gcagcagcaa ga   22

<210> 120
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<212> DNA
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<220>
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<400> 120
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<210> 121
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<400> 121
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<210> 122
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<400> 122
gagcatgatc ctctgcttct c   21

<210> 123
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<400> 123
tctgaagaga agcagaggat ca   22

<210> 124
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<220>
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<400> 124
cagtctttga tttaaccgct tg   22

<210> 125
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<212> DNA
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<220>
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<400> 125
cattgcagtt gaaagaagaa gaaa   24

<210> 126
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<220>
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<400> 126
tgccttgaag agactgagga c   21

<210> 127
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<212> DNA
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<220>
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<400> 127
tctcttcaag gcattctgaa aa   22

<210> 128
<211> 20
<212> DNA
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<400>  128
atccagggca ggaagagact                                                    20


<210>  129
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<212>  DNA
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agcggcacga caattatgta                                                    20


<210>  130
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<220>
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<400>  130
ttttccatta ctttcccaag ga                                                 22


<210>  131
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<212>  DNA
<213>  Artificial

<220>
<223>  artificially synthesized primer seuqnece called MINT23-S

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ctgggtcctt gggaaagtaa                                                    20


<210>  132
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<220>
<223>  artificially synthesized primer seuqnece called MINT23-AS

<400>  132
cagcactgga ggacaaatca                                                    20
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